

SEQUENCE LISTING

<110> Bertin, John

<120> NOVEL MOLECULES OF THE CARD-RELATED PROTEIN FAMILY AND USES THEREOF

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<141> 2001-01-22

<150> 60/181,159

<151> 2000-02-09

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Leu	Āsp	Leu	Leu	Lys	Thr	Arq	Glv	Lvs	Asn	Glv	Ala	Tle	Ala	Phe	Len	4/3
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	20	00 1100 OIU	25	3:		
	_~			J	~	
tgc atc tg	c ccc agc c	gc ctc acc	ccc tac ctg	cac caa ac	c aaq ata	144
			Pro Tyr Leu			
- 4		-	2 - · ·		4	

45

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	agg Arg														864
	aag Lys 290						-	-	_	-		_	_		912
	aag Lys														960
	gag Glu														1008
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	gtc Val														1104
	ctg Leu 370														1152
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	agc Ser					-			-		_			_	1296
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	gac Asp	_	_	_	_		_	_				_		_	1488

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-		_			gaa Glu	_					_	_	_	_		1632
				-	ccg Pro 550		-	_	_		_	_	_	~ ~	_	1680
		_	_		gag Glu			-	_	_	_		_	_	-	1728
-	_	-	_	_	cca Pro	-						_	_	_		1776
_	-		_		cgt Arg		_	-	_	_		-		_		1824
			_	_	agg Arg		_	_	_	-		-	_			1872
					ctg Leu 630											1920
					cac His				_		_			-	_	1968
					ggc Gly											2016
					aag Lys											2064
					agg Arg											2112
					ggt Gly 710											2160
aaa	gtg	gcg	acc	tcg	ggg	gac	tca	ttc	tac	atc	cgg	gtc	aac	ctg	gcc	2208

Ly	s Val	Ala	a Thr	Ser 725	Gly	Asp	Ser	· Phe	Tyr 730		e Arg	y Val	Asn	Leu 735		
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ct Le	g cac u His	gto Val 755	. Thr	gac Asp	acc Thr	atg Met	ttc Phe 760	Gln	ggc Gly	tgc Cys	ggc	tgc Cys 765	tgg Trp	cat His	gcc Ala	2304
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ga As _l	c atg p Met	act Thr	cag Gln	cag Gln 805	Cys	acc Thr	gtg Val	acc Thr	cgc Arg 810	aag Lys	cca Pro	tct Ser	tct Ser	ggg Gly 815	gga Gly	2448
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The state of the s	agc Ser	cag Gln	agg Arg	acg Thr 1060	Gly	atc Ile	gcc Ala	acc Thr	cag Gln 106	Gln	cgc Arg	cag Gln	tgt Cys	cac His	cga Arg)	aga Arg	3216
1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1	att Ile	aac Asn	cca Pro 1075	Arg	cag Gln	agg Arg	atg Met	ggc Gly 1080	Ile	gcc Ala	acc Thr	cag Gln	caa Gln 1085	Arg	cag Gln	tgt Cys	3264
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Asp Ser Asp Cys Ser Leu Val Ser Ser Thr Glu Ser Gln Leu Leu Ser Asp Leu Ser Ala Thr Ser Ser Arg Glu Leu Val Asp Ser Phe Arg Ser Ser Ser Pro Ala Pro Pro Ser Gln Gln Ser Leu Tyr Lys Arg Val Ala Glu Asp Phe Gly Glu Glu Pro Trp Ser Phe Ser Ser Cys Leu Glu Ile Pro Glu Gly Asp Pro Gly Ala Leu Pro Gly Ala Lys Ala Gly Asp Pro His Leu Asp Tyr Glu Leu Leu Asp Thr Ala Asp Leu Pro Gln Leu Glu Ser Ser Leu Gln Pro Val Ser Pro Gly Arg Leu Asp Val Ser Glu Ser Ala Gln Ala Gly Arg Leu Pro Ala Cys Ser Gly Val Leu Met Arg Arg Arg Pro Ala Arg Arg Ile Leu Ser Gln Val Thr Met Leu Ala Phe Gln Gly Asp Ala Leu Leu Glu Gln Ile Ser Val Ile Gly Gly Asn Leu Thr Gly Ile Phe Ile His Arg Val Thr Pro Gly Ser Ala Ala Asp Gln Met Ala Leu Arg Pro Gly Thr Gln Ile Val Met Val Asp Tyr Glu Ala Ser Glu Pro Leu Phe Lys Ala Val Leu Glu Asp Thr Thr Leu Glu Glu Ala Val Gly Leu Leu Arg Arg Val Asp Gly Phe Cys Cys Leu Ser Val Lys Val Asn Thr Asp Gly Tyr Lys Arg Leu Leu Gln Asp Leu Glu Ala Lys Val Ala Thr Ser Gly Asp Ser Phe Tyr Ile Arg Val Asn Leu Ala Met Glu Gly Arg Ala Lys Gly Glu Leu Gln Val His Cys Asn Glu Val Leu His Val Thr Asp Thr Met Phe Gln Gly Cys Gly Cys Trp His Ala His Arg Val Asn Ser Tyr Thr Met Lys Asp Thr Ala Ala His Gly Thr Ile Pro Asn Tyr Ser Arg Ala Gln Gln Leu Ile Ala Leu Ile Gln Asp Met Thr Gln Gln Cys Thr Val Thr Arg Lys Pro Ser Ser Gly Gly Pro Gln Lys Leu Val Arg Ile Val Ser Met Asp Lys Ala Lys Ala Ser Pro Leu Arg Leu Ser Phe Asp Arg Gly Gln Leu Asp Pro Ser Arg Met Glu Gly Ser Ser Thr Cys Phe Trp Ala Glu Ser Cys Leu Thr Leu Val Pro Tyr Thr Leu Val Arg Pro His Arg Pro Ala Arg Pro Arg Pro Val Leu Leu Val Pro Arg Ala Val Gly Lys Ile Leu Ser Glu Lys Leu Cys Leu 890 895 Leu Gln Gly Phe Lys Lys Cys Leu Ala Glu Tyr Leu Ser Gln Glu Glu Tyr Glu Ala Trp Ser Gln Arg Gly Asp Ile Ile Gln Glu Gly Glu Val

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                                    955
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              965
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Gln Arg Thr Gly Ile Ala Thr Gln Gln Arg Gln Cys His Arg Arg Ile
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Arg Arg Ile Asn Pro Ser Gln Arg Thr Gly Ile Thr Thr Gln Gln Cys
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   1090
                                         1100
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                          40
Arg Arg Asp Lys Ala Arg Glu Leu Ile Asp Ser Val Gln Lys Lys Gly
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Leu Arg Asp
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Glu Thr Val Arg Gln Val Ala Lys Gln Gly Lys Ile Cys Ile Leu Asp
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Val Glu Pro Gln Gly Val Lys Arg Leu Arg Thr Ala Glu Leu Ser Asn
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